RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/501, 841
Source:	PLT.
Date Processed by STIC:	12/08/2005

ENTERED



PCT

RAW SEQUENCE LISTING DATE: 12/08/2005
PATENT APPLICATION: US/10/501,841 TIME: 10:26:47

Input Set : A:\-144-1-3.app

```
3 <110> APPLICANT: Gaiger, Alexander
        Algate, Paul A.
 5
        Mannion, Jane
                                                           (pg-6)
        Clapper, Jonathan David
 6
 7
        Wang, Aijun
 8
        Ordonez, Nadia
 9
        Carter, Lauren
10
        McNeill, Patricia Dianne
11
        Corixa Corporation
13 <120> TITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis
        and Therapy of Hematological Malignancies
16 <130> FILE REFERENCE: 014058-014402PC
18 <140> CURRENT APPLICATION NUMBER: US 10/501,841
19 <141> CURRENT FILING DATE: 2004-07-14
21 <150> PRIOR APPLICATION NUMBER: US 10/057,475
22 <151> PRIOR FILING DATE: 2002-01-22
24 <150> PRIOR APPLICATION NUMBER: WO PCT/US03/02353
25 <151> PRIOR FILING DATE: 2003-01-22
27 <160> NUMBER OF SEQ ID NOS: 124
29 <170> SOFTWARE: PatentIn Ver. 2.1
31 <210> SEO ID NO: 1
32 <211> LENGTH: 2672
33 <212> TYPE: DNA
34 <213> ORGANISM: Homo sapiens
36 <400> SEQUENCE: 1
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39 gccgtgactt tccccctgaa gtccaaagta aagcaagttg actctattgt ctggaccttc 180
40 aacacaaccc ctcttgtcac catacagcca gaagggggca ctatcatagt gacccaaaat 240
41 cgtaataggg agagagtaga cttcccagat ggaggctact ccctgaagct cagcaaactg 300
42 aagaagaatg actcagggat ctactatgtg gggatataca gctcatcact ccagcagccc 360
43 tecacecagg agtacgtget geatgtetae gageacetgt caaageetaa agteaceatg 420
44 ggtctgcaga gcaataagaa tggcacctgt gtgaccaatc tgacatgctg catggaacat 480
45 ggggaagagg atgtgattta tacctggaag gccctggggc aagcagccaa tgagtcccat 540
46 aatgggteca tectececat etectggaga tggggagaaa gtgatatgae etteatetge 600
47 gttgccagga accetgtcag cagaaaette teaageeeca teettgecag gaagetetgt 660
48 gaaggtgetg etgatgacce agatteetee atggteetee tgtgteteet gttggtgeee 720
49 ctcctgctca gtctctttgt actggggcta tttctttggt ttctgaagag agagagacaa 780
50 gaagagtaca ttgaagagaa gaagagagtg gacatttgtc gggaaactcc taacatatgc 840
51 ccccattctg gagagaacac agagtacgac acaatccctc acactaatag aacaatccta 900
52 aaggaagatc cagcaaatac ggtttactcc actgtggaaa taccgaaaaa gatggaaaat 960
53 ccccactcac tgctcacgat gccagacaca ccaaggctat ttgcctatga gaatgttatc 1020
54 tagacagcag tgcactcccc taagtctctg ctcaaaaaaa aaacaattct cggcccaaag 1080
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Input Set : A:\-144-1-3.app

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56 gacttttttc caggataaat tatctctgat gcttctttag atttaagagt tcataattcc 1200
57 atccactgct gagaaatctc ctcaaaccca gaaggtttaa tcacttcatc ccaaaaatgg 1260
58 gattgtgaat gtcagcaaac cataaaaaaa gtgcttagaa gtattcctat agaaatgtaa 1320
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60 ctggagtttc attccatccc agggcttgga tgtaaggatt ataccaagag tcttgctacc 1440
61 aggagggcaa gaagaccaaa acagacagac aagtccagca gaagcagatg cacctgacaa 1500
62 aaatggatgt attaattggc tctataaact atgtgcccag cactatgctg agcttacact 1560
63 aattggtcag acgtgctgtc tgccctcatg aaattggctc caaatgaatg aactactttc 1620
64 atgagcagtt gtagcaggcc tgaccacaga ttcccagagg gccaggtgtg gatccacagg 1680
65 acttgaaggt caaagttcac aaagatgaag aatcagggta gctgaccatg tttggcagat 1740
66 actataatgg agacacagaa gtgtgcatgg cccaaggaca aggacctcca gccaggcttc 1800
67 atttatgcac ttgtgctgca aaagaaaagt ctaggtttta aggctgtgcc agaacccatc 1860
68 ccaataaaga gaccgagtct gaagtcacat tgtaaatcta gtgtaggaga cttggagtca 1920
69 ggcagtgaga ctggtggggc acggggggca gtgggtactt gtaaaccttt aaagatggtt 1980
70 aattcattca atagatattt attaagaacc tatgcggccc ggcatggtgg ctcacacctg 2040
71 taatcccagc actttgggag gccaaggtgg gtgggtcatc tgaggtcagg agttcaagac 2100
72 cagcetggee aacatggtga aaccecatet etactaaaga tacaaaaatt tgetgagegt 2160
73 ggtggtgtgc acctgtaatc ccagctactc gagaggccaa ggcatgagaa tcgcttgaac 2220
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76 aagatgaaca teeetaecaa cacagagete accatetett ataettaagt gaaaaacatg 2400
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78 gacctcccta ccaagtgatg aaagtgttga aaaacttaat aacaaatgct tgttgggcaa 2520
79 gaatgggatt gaggattatc ttctctcaga aaggcattgt gaaggaattg agccagatct 2580
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81 agatattgtg agattcaaaa aaaaaaaaaa aa
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85 <211> LENGTH: 335
86 <212> TYPE: PRT
87 <213 > ORGANISM: Homo sapiens
89 <400> SEQUENCE: 2
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                     5
93 Leu Thr Gly Ser Ala Ala Ser Gly Pro Val Lys Glu Leu Val Gly Ser
                                    25
96 Val Gly Gly Ala Val Thr Phe Pro Leu Lys Ser Lys Val Lys Gln Val
                                40
99 Asp Ser Ile Val Trp Thr Phe Asn Thr Thr Pro Leu Val Thr Ile Gln
                             55
102 Pro Glu Gly Gly Thr Ile Ile Val Thr Gln Asn Arg Asn Arg Glu Arg
103
                         70
                                             75
105 Val Asp Phe Pro Asp Gly Gly Tyr Ser Leu Lys Leu Ser Lys Leu Lys
                     85
                                         90
108 Lys Asn Asp Ser Gly Ile Tyr Tyr Val Gly Ile Tyr Ser Ser Leu
                100
                                    105
111 Gln Gln Pro Ser Thr Gln Glu Tyr Val Leu His Val Tyr Glu His Leu
                                120
114 Ser Lys Pro Lys Val Thr Met Gly Leu Gln Ser Asn Lys Asn Gly Thr
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Input Set : A:\-144-1-3.app

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115
        130
                            135
117 Cys Val Thr Asn Leu Thr Cys Cys Met Glu His Gly Glu Glu Asp Val
                        150
120 Ile Tyr Thr Trp Lys Ala Leu Gly Gln Ala Ala Asn Glu Ser His Asn
                    165
                                         170
123 Gly Ser Ile Leu Pro Ile Ser Trp Arg Trp Gly Glu Ser Asp Met Thr
124
                180
                                    185
126 Phe Ile Cys Val Ala Arg Asn Pro Val Ser Arg Asn Phe Ser Ser Pro
            195
                                200
                                                     205
129 Ile Leu Ala Arg Lys Leu Cys Glu Gly Ala Ala Asp Asp Pro Asp Ser
                            215
132 Ser Met Val Leu Leu Cys Leu Leu Leu Val Pro Leu Leu Ser Leu
133 225
                        230
                                            235
135 Phe Val Leu Gly Leu Phe Leu Trp Phe Leu Lys Arg Glu Arg Gln Glu
138 Glu Tyr Ile Glu Glu Lys Lys Arg Val Asp Ile Cys Arg Glu Thr Pro
139
                                    265
141 Asn Ile Cys Pro His Ser Gly Glu Asn Thr Glu Tyr Asp Thr Ile Pro
                                280
144 His Thr Asn Arg Thr Ile Leu Lys Glu Asp Pro Ala Asn Thr Val Tyr
145
        290
                            295
147 Ser Thr Val Glu Ile Pro Lys Lys Met Glu Asn Pro His Ser Leu Leu
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                                            315
150 Thr Met Pro Asp Thr Pro Arg Leu Phe Ala Tyr Glu Asn Val Ile
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155 <211> LENGTH: 834
156 <212> TYPE: DNA
157 <213> ORGANISM: Homo sapiens
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162 catgettgea tacettgtea aettegatgt tettetaata etecteetet aacatgteag 180
163 cgttattgta atgcaagtgt gaccaattca gtgaaaggaa cgaatgcgat tctctggacc 240
164 tgtttgggac tgagcttaat aatttctttg gcagttttcg tgctaatgtt tttgctaagg 300
165 aagataagct ctgaaccatt aaaggacgag tttaaaaaca caggatcagg tctcctgggc 360
166 atggctaaca ttgacctgga aaagagcagg actggtgatg aaattattct tccqagaggc 420
167 ctcgagtaca cggtggaaga atgcacctgt gaagactgca tcaagagcaa accgaaggtc 480
168 gactetgace attgetttee acteceaget atggaggaag gegeaaceat tettgteace 540
169 acgaaaacga atgactattg caagagcctg ccagctgctt tgagtgctac ggagatagag 600
170 aaatcaattt ctgctaggta attaaccatt tcgactcgag cagtgccact ttaaaaatct 660
171 tttgtcagaa tagatgatgt gtcagatctc tttaggatga ctgtattttt cagttgccga 720
172 tacagetttt tgteetetaa etgtggaaac tetttatgtt agatatattt etetaggtta 780
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179 <213> ORGANISM: Homo sapiens
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Input Set : A:\-144-1-3.app

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185 Leu Leu His Ala Cys Ile Pro Cys Gln Leu Arg Cys Ser Ser Asn Thr
186
                 20
188 Pro Pro Leu Thr Cys Gln Arg Tyr Cys Asn Ala Ser Val Thr Asn Ser
189
                                 40
191 Val Lys Gly Thr Asn Ala Ile Leu Trp Thr Cys Leu Gly Leu Ser Leu
194 Ile Ile Ser Leu Ala Val Phe Val Leu Met Phe Leu Leu Arg Lys Ile
195
                         70
197 Ser Ser Glu Pro Leu Lys Asp Glu Phe Lys Asn Thr Gly Ser Gly Leu
                                         90
200 Leu Gly Met Ala Asn Ile Asp Leu Glu Lys Ser Arg Thr Gly Asp Glu
                100
                                    105
203 Ile Ile Leu Pro Arg Gly Leu Glu Tyr Thr Val Glu Glu Cys Thr Cys
                                120
206 Glu Asp Cys Ile Lys Ser Lys Pro Lys Val Asp Ser Asp His Cys Phe
207
        130
                            135
209 Pro Leu Pro Ala Met Glu Glu Gly Ala Thr Ile Leu Val Thr Thr Lys
210 145
                        150
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212 Thr Asn Asp Tyr Cys Lys Ser Leu Pro Ala Ala Leu Ser Ala Thr Glu
213
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                                        170
215 Ile Glu Lys Ser Ile Ser Ala Arg
216
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221 <212> TYPE: DNA
222 <213> ORGANISM: Homo sapiens
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226 gggggccctg aggatcctcc cagaagtaaa ggtagagggg gagctgggcg gatcagttac 120
227 catcaagtgc ccacttcctg aaatgcatgt gaggatatat ctgtgccggg agatggctgg 180
228 atctggaaca tgtggtaccg tggtatccac caccaacttc atcaaggcag aatacaaggg 240
229 ccgagttact ctgaagcaat acccacgcaa gaatctgttc ctagtggagg taacacagct 300
230 gacagaaagt gacageggag tetatgeetg eggageggge atgaacacag aceggggaaa 360
231 gacccagaaa gtcaccctga atgtccacag tgaatacgag ccatcatggg aagagcagcc 420
232 aatgcctgag actccaaaat ggtttcatct gccctatttg ttccagatgc ctgcatatgc 480
233 cagttettee aaattegtaa eeagagttae cacaceaget caaaggggea aggteeetee 540
234 agttcaccac tcctccccca ccacccaaat cacccaccgc cctcgagtgt ccagagcatc 600
235 ttcagtagca ggtgacaagc cccgaacctt cctgccatcc actacagcct caaaaatctc 660
236 agetetggag gggetgetea ageeceagae geecagetae aaceaceaea eeaggetgea 720
237 caggcagaga gcactggact atggctcaca gtctgggagg gaaggccaag gatttcacat 780
238 cctgatcccg accatcctgg gccttttcct gctggcactt ctggggctgg tggtgaaaag 840
239 ggccgttgaa aggaggaaag ccctctccag gcgggcccgc cgactggccg tgaggatgcg 900
240 cgccctggag agctcccaga ggccccgcgg gtcgccgcga ccgcgctccc aaaacaacat 960
241 ctacagegec tgccegegge gegetegtgg ageggaeget geaggeacag gggaageece 1020
242 cgttcccggc cccggagcgc cgttgccccc cgccccgctg caggtgtctg aatctccctg 1080
243 getecatgee ceatetetga agaccagetg tgaataegtg ageetetace accageetge 1140
244 cgccatgatg gaggacagtg attcagatga ctacatcaat gttcctgcct gacaactccc 1200
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Input Set : A:\-144-1-3.app

Output Set: N:\CRF4\12082005\J501841.raw

245 cagetatece ecaaceecag geteggactg tggtgecaag gagteteate tatetgetga 1260 246 tgtccaatac ctgcttcatg tgttctcaga gccctcatca ttcccatgcc ccatctcgat 1320 247 cccatcccca tctatctgt 250 <210> SEQ ID NO: 6 251 <211> LENGTH: 390 252 <212> TYPE: PRT 253 <213> ORGANISM: Homo sapiens 255 <400> SEQUENCE: 6 256 Met Asp Phe Trp Leu Trp Pro Leu Tyr Phe Leu Pro Val Ser Gly Ala 259 Leu Arg Ile Leu Pro Glu Val Lys Val Glu Gly Glu Leu Gly Gly Ser 25 262 Val Thr Ile Lys Cys Pro Leu Pro Glu Met His Val Arg Ile Tyr Leu 35 40 265 Cys Arg Glu Met Ala Gly Ser Gly Thr Cys Gly Thr Val Val Ser Thr 268 Thr Asn Phe Ile Lys Ala Glu Tyr Lys Gly Arg Val Thr Leu Lys Gln 269 65 271 Tyr Pro Arg Lys Asn Leu Phe Leu Val Glu Val Thr Gln Leu Thr Glu 85 90 274 Ser Asp Ser Gly Val Tyr Ala Cys Gly Ala Gly Met Asn Thr Asp Arg 100 105 277 Gly Lys Thr Gln Lys Val Thr Leu Asn Val His Ser Glu Tyr Glu Pro 115 120 125 280 Ser Trp Glu Glu Gln Pro Met Pro Glu Thr Pro Lys Trp Phe His Leu 135 283 Pro Tyr Leu Phe Gln Met Pro Ala Tyr Ala Ser Ser Ser Lys Phe Val 284 145 150 155 286 Thr Arg Val Thr Thr Pro Ala Gln Arg Gly Lys Val Pro Pro Val His 289 His Ser Ser Pro Thr Thr Gln Ile Thr His Arg Pro Arg Val Ser Arg 180 185 292 Ala Ser Ser Val Ala Gly Asp Lys Pro Arg Thr Phe Leu Pro Ser Thr 200 295 Thr Ala Ser Lys Ile Ser Ala Leu Glu Gly Leu Leu Lys Pro Gln Thr 215 298 Pro Ser Tyr Asn His His Thr Arg Leu His Arg Gln Arg Ala Leu Asp 230 235 301 Tyr Gly Ser Gln Ser Gly Arg Glu Gly Gln Gly Phe His Ile Leu Ile 245 250 304 Pro Thr Ile Leu Gly Leu Phe Leu Leu Ala Leu Leu Gly Leu Val Val 260 265 307 Lys Arg Ala Val Glu Arg Arg Lys Ala Leu Ser Arg Arg Ala Arg Arg 275 280 310 Leu Ala Val Arg Met Arg Ala Leu Glu Ser Ser Gln Arg Pro Arg Gly 295 300 313 Ser Pro Arg Pro Arg Ser Gln Asn Asn Ile Tyr Ser Ala Cys Pro Arg 316 Arg Ala Arg Gly Ala Asp Ala Ala Gly Thr Gly Glu Ala Pro Val Pro

Input Set : A:\-144-1-3.app

Output Set: N:\CRF4\12082005\J501841.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:16; N Pos. 9 Seq#:22; N Pos. 971,975,997,1060 Seg#:30; N Pos. 488,518 Seq#:36; N Pos. 488,518 Seq#:37; Xaa Pos. 5,82 Seq#:38; N Pos. 340,529,534,547 Seq#:41; N Pos. 13,50,82,85,100 Seq#:49; N Pos. 152,174,180 Seq#:50; Xaa Pos. 50,58,60 Seq#:51; N Pos. 152,174,180 Seq#:52; Xaa Pos. 50,58,60 Seq#:67; N Pos. 16 Seq#:72; N Pos. 519 Seq#:78; N Pos. 150,183,523 Seq#:90; N Pos. 525,533 Seq#:96; N Pos. 1619,1628 Seq#:98; N Pos. 271,283,311 Seq#:105; N Pos. 19,67,109 Seq#:110; N Pos. 467

Seq#:112; N Pos. 90,137,161

DATE: 12/08/2005

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/501,841 TIME: 10:26:48

Input Set : A:\-144-1-3.app

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L:1195 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:1743 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:960
L:1744 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:1020
L:2147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:480
L:2524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:480
L:2539 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0
L:2554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:80
L:2574 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:300
L:2577 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:480
L:2578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:540
L:2702 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0
L:2703 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:60
L:3184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 after pos.:120
L:3208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:48
L:3225 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:120
L:3249 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:48
L:3932 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:0
L:4077 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72 after pos.:480
L:4323 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:120
L:4324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:180
L:4329 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:480
L:5360 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:90 after pos.:480
L:5627 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:96 after pos.:1560
L:5628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:96 after pos.:1620
L:5694 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:240
L:5695 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:300
L:5919 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:105 after pos.:0
L:5920 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:105 after pos.:60
L:6237 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:110 after pos.:420
L:6270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 after pos.:60
L:6271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 after pos.:120
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